

PCT09

RAW SEQUENCE LISTING DATE: 08/14/2001  
 PATENT APPLICATION: US/09/890,688 TIME: 11:01:23

Input Set : A:\Sequence Listing Kato et al.txt  
 Output Set: N:\CRF3\08142001\I890688.raw

3 <110> APPLICANT: Seishi KATO  
 4 Chikashi EGUCHI  
 5 Mihoro SAEKI  
 7 <120> TITLE OF INVENTION: Human Proteins and cDNAs thereof  
 9 <130> FILE REFERENCE: 2001-1102A/WMC/00653  
 C--> 11 <140> CURRENT APPLICATION NUMBER: US/09/890,688  
 C--> 12 <141> CURRENT FILING DATE: 2001-08-06  
 14 <150> PRIOR APPLICATION NUMBER: JP 11-346863  
 15 <151> PRIOR FILING DATE: 1999-12-06  
 17 <150> PRIOR APPLICATION NUMBER: JP 11-34684  
 18 <151> PRIOR FILING DATE: 1999-12-06  
 20 <150> PRIOR APPLICATION NUMBER: JP 2000-31062  
 21 <151> PRIOR FILING DATE: 2000-02-08  
 23 <150> PRIOR APPLICATION NUMBER: JP 2000-34091  
 24 <151> PRIOR FILING DATE: 2000-02-10  
 26 <150> PRIOR APPLICATION NUMBER: JP 2000-34090  
 27 <151> PRIOR FILING DATE: 2000-02-10  
 29 <150> PRIOR APPLICATION NUMBER: JP 2000-35829  
 30 <151> PRIOR FILING DATE: 2000-02-14  
 32 <150> PRIOR APPLICATION NUMBER: JP 2000-35899  
 33 <151> PRIOR FILING DATE: 2000-02-14  
 35 <150> PRIOR APPLICATION NUMBER: JP 2000-71161  
 36 <151> PRIOR FILING DATE: 2000-03-14  
 38 <150> PRIOR APPLICATION NUMBER: JP 2000-160851  
 39 <151> PRIOR FILING DATE: 2000-05-30  
 41 <160> NUMBER OF SEQ ID NOS: 160  
 43 <170> SOFTWARE: PatentIn Ver. 2.1  
 45 <210> SEQ ID NO: 1  
 46 <211> LENGTH: 1323  
 47 <212> TYPE: DNA  
 48 <213> ORGANISM: Homo sapiens  
 50 <220> FEATURE:  
 51 <221> NAME/KEY: CDS  
 52 <222> LOCATION: (27)..(881)  
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 59 gcg gga aga ctc ttt gaa atg cct gcg gtg cta gag cga ctg agc cgc 101  
 60 Ala Gly Arg Leu Phe Glu Met Pro Ala Val Leu Glu Arg Leu Ser Arg  
 61 10 15 20 25  
 63 tat aat agc acg tcc caa gct ttt gct gag gtg ctg cgg ctg ccg aag 149  
 64 Tyr Asn Ser Thr Ser Gln Ala Phe Ala Glu Val Leu Arg Leu Pro Lys  
 65 30 35 40  
 67 cag cag ctg agg aag ctg ctg tac ccg ctg cag gaa gta gag cgg ttc 197  
 68 Gln Gln Leu Arg Lys Leu Tyr Pro Leu Gln Glu Val Glu Arg Phe  
 69 45 50 55

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71 ctc gcc ccc tac ggg agg caa gac ctt cac ctg cgt atc ttt gac cca	245
72 Leu Ala Pro Tyr Gly Arg Gln Asp Leu His Leu Arg Ile Phe Asp Pro	
73       60                         65                         70	
75 agc ccg gag gac ata gcc agg gcg gac aac atc ttc acg gcc act gaa	293
76 Ser Pro Glu Asp Ile Ala Arg Ala Asp Asn Ile Phe Thr Ala Thr Glu	
77       75                         80                         85	
79 cgg aac cgc atc gac tac gtc agc tcc gcc gtc cgt atc gac cac gcc	341
80 Arg Asn Arg Ile Asp Tyr Val Ser Ser Ala Val Arg Ile Asp His Ala	
81       90                         95                         100                         105	
83 ccg gac ctt ccg cgg cca gag gtg tgt ttt ata ggc aga agc aat gtt	389
84 Pro Asp Leu Pro Arg Pro Glu Val Cys Phe Ile Gly Arg Ser Asn Val	
85       110                         115                         120	
87 gga aaa tca tct cta atc aag gct tta ttt tca ctg gcc cct gag gtt	437
88 Gly Lys Ser Ser Leu Ile Lys Ala Leu Phe Ser Leu Ala Pro Glu Val	
89       125                         130                         135	
91 gaa gtc aga gtc tcc aaa aaa cca gga cac aca aag aaa atg aat ttt	485
92 Glu Val Arg Val Ser Lys Lys Pro Gly His Thr Lys Lys Met Asn Phe	
93       140                         145                         150	
95 ttc aaa gtt gga aaa cat ttt aca gtg gtg gac atg cca ggt tat ggc	533
96 Phe Lys Val Gly Lys His Phe Thr Val Val Asp Met Pro Gly Tyr Gly	
97       155                         160                         165	
99 ttt aga gca cct gaa gat ttt gtt gac atg gta gag acc tat cta aaa	581
100 Phe Arg Ala Pro Glu Asp Phe Val Asp Met Val Glu Thr Tyr Leu Lys	
101 170                             175                         180                                 185	
103 gaa cga agg aac ttg aag aga aca ttt tta tta gtg gat agc gtt gtt	629
104 Glu Arg Arg Asn Leu Lys Arg Thr Phe Leu Leu Val Asp Ser Val Val	
105       190                         195                         200	
107 gga att caa aaa aca gac aat att gcc ata gaa atg tgt gaa gaa ttt	677
108 Gly Ile Gln Lys Thr Asp Asn Ile Ala Ile Glu Met Cys Glu Glu Phe	
109       205                         210                         215	
111 gca tta cct tat gtg att gta tta aca aaa att gac aaa tct tcc aag	725
112 Ala Leu Pro Tyr Val Ile Val Leu Thr Lys Ile Asp Lys Ser Ser Lys	
113       220                         225                         230	
115 gga cat ctt tta aaa caa gtg ctt cag atc cag aaa ttt gtt aac atg	773
116 Gly His Leu Leu Lys Gln Val Leu Gln Ile Gln Lys Phe Val Asn Met	
117       235                         240                         245	
119 aaa act caa gga tgt ttt cct cag ttg ttt cct gta agt gct gtg acc	821
120 Lys Thr Gln Gly Cys Phe Pro Gln Leu Phe Pro Val Ser Ala Val Thr	
121 250                             255                         260                                 265	
123 ttt tct gga atc cac ctg ttg aga tgc ttt ata gcc agt gta aca gga	869
124 Phe Ser Gly Ile His Leu Leu Arg Cys Phe Ile Ala Ser Val Thr Gly	
125       270                         275                         280	
127 agt ctt gac taa tggtccccgg ttttagctgaa gattcaaaaag ctttatgcta	921
128 Ser Leu Asp	
131 actggaggtta aatacctaga agaatttcaa cattgtttta aatgttgtgc atctgttaact	981
133 tcaggaggat cacttgagct taaaaacctg tgccttctcg aaacaagaat ttgtgcctga	1041
135 ggtaaaaaaa gttgtaaagt tattgaatta tggtgttcat tagaacagct actagctgat	1101
137 tcccctattt taacaaactg acaagagcac atccataaaa tgaaaacctg ttacaactat	1161
139 gtacagaagg gtttgacggtt ttattgggct ttatgtttt aaagaatatg tctactatgg	1221

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Input Set : A:\Sequence Listing Kato et al.txt  
 Output Set: N:\CRF3\08142001\I890688.raw

141 gtatTTTT tttaaatgtt aaaatggct aggtaaaagg gggctgctt tctgttaagc 1281  
 143 atcgataggt aagttagtgg ataaaagtta ctatgttaagc ct 1323  
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 148 <212> TYPE: PRT  
 149 <213> ORGANISM: Homo sapiens  
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 153 1 5 10 15  
 155 Pro Ala Val Leu Glu Arg Leu Ser Arg Tyr Asn Ser Thr Ser Gln Ala  
 156 20 25 30  
 158 Phe Ala Glu Val Leu Arg Leu Pro Lys Gln Gln Leu Arg Lys Leu Leu  
 159 35 40 45  
 161 Tyr Pro Leu Gln Glu Val Glu Arg Phe Leu Ala Pro Tyr Gly Arg Gln  
 162 50 55 60  
 164 Asp Leu His Leu Arg Ile Phe Asp Pro Ser Pro Glu Asp Ile Ala Arg  
 165 65 70 75 80  
 167 Ala Asp Asn Ile Phe Thr Ala Thr Glu Arg Asn Arg Ile Asp Tyr Val  
 168 85 90 95  
 170 Ser Ser Ala Val Arg Ile Asp His Ala Pro Asp Leu Pro Arg Pro Glu  
 171 100 105 110  
 173 Val Cys Phe Ile Gly Arg Ser Asn Val Gly Lys Ser Ser Leu Ile Lys  
 174 115 120 125  
 176 Ala Leu Phe Ser Leu Ala Pro Glu Val Glu Val Arg Val Ser Lys Lys  
 177 130 135 140  
 179 Pro Gly His Thr Lys Lys Met Asn Phe Phe Lys Val Gly Lys His Phe  
 180 145 150 155 160  
 182 Thr Val Val Asp Met Pro Gly Tyr Gly Phe Arg Ala Pro Glu Asp Phe  
 183 165 170 175  
 185 Val Asp Met Val Glu Thr Tyr Leu Lys Glu Arg Arg Asn Leu Lys Arg  
 186 180 185 190  
 188 Thr Phe Leu Leu Val Asp Ser Val Val Gly Ile Gln Lys Thr Asp Asn  
 189 195 200 205  
 191 Ile Ala Ile Glu Met Cys Glu Glu Phe Ala Leu Pro Tyr Val Ile Val  
 192 210 215 220  
 194 Leu Thr Lys Ile Asp Lys Ser Ser Lys Gly His Leu Leu Lys Gln Val  
 195 225 230 235 240  
 197 Leu Gln Ile Gln Lys Phe Val Asn Met Lys Thr Gln Gly Cys Phe Pro  
 198 245 250 255  
 200 Gln Leu Phe Pro Val Ser Ala Val Thr Phe Ser Gly Ile His Leu Leu  
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 208 <211> LENGTH: 1120  
 209 <212> TYPE: DNA  
 210 <213> ORGANISM: Homo sapiens  
 212 <220> FEATURE:  
 213 <221> NAME/KEY: CDS

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Input Set : A:\Sequence Listing Kato et al.txt  
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214 <222> LOCATION: (18)..(719)  
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 221 gct ctg ctg cgg gcg gtc gct gga cgg ctt cgg gga ggc gtc cag 98  
 222 Ala Leu Leu Arg Ala Gly Arg Leu Leu Arg Gly Gly Val Gln  
 223 15 20 25  
 225 gag cta ctg cgg ccg cga cat gaa ggg aac gcc cct gac ctg gcc tgc 146  
 226 Glu Leu Leu Arg Pro Arg His Glu Gly Asn Ala Pro Asp Leu Ala Cys  
 227 30 35 40  
 229 aac ttc agc ctt tct caa aat cgg ggc acg gtc atc gtg gag cgc tgg 194  
 230 Asn Phe Ser Leu Ser Gln Asn Arg Gly Thr Val Ile Val Glu Arg Trp  
 231 45 50 55,  
 233 tgg aag gta ccg ctg gcc ggg ggg ggc cgg aag ccg cgc ctg cac cgg 242  
 234 Trp Lys Val Pro Leu Ala Gly Gly Arg Lys Pro Arg Leu His Arg  
 235 60 65 70 75  
 237 cga cat cgc gtc tat aag ctg gtg gag gac acg aag cat cgg ccc aaa 290  
 238 Arg His Arg Val Tyr Lys Leu Val Glu Asp Thr Lys His Arg Pro Lys  
 239 80 85 90  
 241 gaa aac ctg gag ctc atc ctg acg cag tcg gtg gag aat gtt gga gtc 338  
 242 Glu Asn Leu Glu Leu Ile Leu Thr Gln Ser Val Glu Asn Val Gly Val  
 243 95 100 105  
 245 cgg ggt gac ctg tca gtg aag aaa tct tta ggc cgg aat cga ctc 386  
 246 Arg Gly Asp Leu Val Ser Val Lys Lys Ser Leu Gly Arg Asn Arg Leu  
 247 110 115 120  
 249 ctt cct cag gga ctg gct gta tat gca tcc cct gaa aac aag aag ctg 434  
 250 Leu Pro Gln Gly Leu Ala Val Tyr Ala Ser Pro Glu Asn Lys Lys Leu  
 251 125 130 135  
 253 ttt gaa gag gag aaa ttg ctg aga caa gaa gga aaa tta gag aag atc 482  
 254 Phe Glu Glu Glu Lys Leu Leu Arg Gln Glu Gly Lys Leu Glu Lys Ile  
 255 140 145 150 155  
 257 cag acc aag gca ggt gag gcg ctt ggt gtt gtg gtt gcc cca cat aca 530  
 258 Gln Thr Lys Ala Gly Glu Ala Leu Gly Val Val Val Ala Pro His Thr  
 259 160 165 170  
 261 tta aag tta cca gca gag cct atc aca cgg tgg ggc gag tat tgg tgt 578  
 262 Leu Lys Leu Pro Ala Glu Pro Ile Thr Arg Trp Gly Glu Tyr Trp Cys  
 263 175 180 185  
 265 gag gtg acg gta aat ggg ctt gat act gtg aga gtg cct atg tct gtc 626  
 266 Glu Val Thr Val Asn Gly Leu Asp Thr Val Arg Val Pro Met Ser Val  
 267 190 195 200  
 269 gtg aac ttt gag aag ccc aag acc aaa aga tat aag tac tgg tta gcc 674  
 270 Val Asn Phe Glu Lys Pro Lys Thr Lys Arg Tyr Lys Tyr Trp Leu Ala  
 271 205 210 215  
 273 cag caa gct gcc aag gct atg gcc ccc acc agc ccc cag atc taa 719  
 274 Gln Gln Ala Ala Lys Ala Met Ala Pro Thr Ser Pro Gln Ile  
 275 220 225 230  
 277 atctactctc cctccaaggc agcaaaggc aatcgggagc agtggagcag aaatgtgcaa 779  
 279 gcaccctgat ctcactccca gctctgacca aatacagaat ttttagagaac atctgaagac 839

RAW SEQUENCE LISTING DATE: 08/14/2001  
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Input Set : A:\Sequence Listing Kato et al.txt  
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281 atcagactgc actgcgtata catgttgaat tcttcatttt tgccatctt aactgtcatc 899  
 283 actggggcag ggaagtccctg ttccagaagt accaggctgt agatttgata agctagatgc 959  
 285 agtagaccga aaccatccaa aacctgtta gcttcttccat ccattggagt ttattggac 1019  
 287 aaacaggaga gccagccatt gtctccagta cttgcctcat tctcatcatc caaactgaac 1079  
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 293 <211> LENGTH: 233  
 294 <212> TYPE: PRT  
 295 <213> ORGANISM: Homo sapiens  
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 302 20 25 30  
 304 Arg His Glu Gly Asn Ala Pro Asp Leu Ala Cys Asn Phe Ser Leu Ser  
 305 35 40 45  
 307 Gln Asn Arg Gly Thr Val Ile Val Glu Arg Trp Trp Lys Val Pro Leu  
 308 50 55 60  
 310 Ala Gly Gly Arg Lys Pro Arg Leu His Arg Arg His Arg Val Tyr  
 311 65 70 75 80  
 313 Lys Leu Val Glu Asp Thr Lys His Arg Pro Lys Glu Asn Leu Glu Leu  
 314 85 90 95  
 316 Ile Leu Thr Gln Ser Val Glu Asn Val Gly Val Arg Gly Asp Leu Val  
 317 100 105 110  
 319 Ser Val Lys Lys Ser Leu Gly Arg Asn Arg Leu Leu Pro Gln Gly Leu  
 320 115 120 125  
 322 Ala Val Tyr Ala Ser Pro Glu Asn Lys Lys Leu Phe Glu Glu Lys  
 323 130 135 140  
 325 Leu Leu Arg Gln Glu Gly Lys Leu Glu Lys Ile Gln Thr Lys Ala Gly  
 326 145 150 155 160  
 328 Glu Ala Leu Gly Val Val Val Ala Pro His Thr Leu Lys Leu Pro Ala  
 329 165 170 175  
 331 Glu Pro Ile Thr Arg Trp Gly Glu Tyr Trp Cys Glu Val Thr Val Asn  
 332 180 185 190  
 334 Gly Leu Asp Thr Val Arg Val Pro Met Ser Val Val Asn Phe Glu Lys  
 335 195 200 205  
 337 Pro Lys Thr Lys Arg Tyr Lys Tyr Trp Leu Ala Gln Gln Ala Ala Lys  
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 351 <222> LOCATION: (90)..(284)  
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**VERIFICATION SUMMARY**

PATENT APPLICATION: US/09/890,688

DATE: 08/14/2001

TIME: 11:01:24

Input Set : A:\Sequence Listing Kato et al.txt  
Output Set: N:\CRF3\08142001\I890688.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application Number  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:6698 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:105  
L:6708 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:105  
L:6735 M:257 W: Feature value mis-spelled or invalid, <221> Name/Key for SEQ ID#:106  
L:6740 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:106